

Table 1

Image Clone ID	Accession No.	Nuc Seq Id	Cluster Id	Cluster Title	Gene	Sensitivity or Resistance
26617	R39862	g797478	Hs.10247	activated leucocyte cell adhesion molecule	ALCAM	Resistance
26617	R13558	g766634	Hs.10247	activated leucocyte cell adhesion molecule	ALCAM	Resistance
27544	R40057	g822754	Hs.112360	prominin (mouse)-like 1	PROML1	Sensitivity
32567	R43511	g821440	Hs.34853	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	ID4	Sensitivity
32567	R20393	g775027	Hs.34853	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	ID4	Sensitivity
33267	R43855	g821734	Hs.162	insulin-like growth factor binding protein 2 (36kD)	IGFBP2	Resistance
51221	H19246	g885486	Hs.106635	ortholog of rat pippin	PIPPIN	Sensitivity
51221	H19245	g885485	Hs.106635	ortholog of rat pippin	PIPPIN	Sensitivity
66317	T66816	g676256	Hs.7644	H1 histone family, member 2	H1F2	Sensitivity
66317	T66815	g676255	Hs.7644	H1 histone family, member 2	H1F2	Sensitivity
66420	R16069	g767878	Hs.189713	ESTs	-	Sensitivity
66498	R16030	g768012	Hs.21688	ESTs	-	Resistance
66498	R16130	g767939	Hs.21688	ESTs	-	Resistance
121661	T97616	g746961	Hs.226410	ESTs	-	Sensitivity
122906	T99784	g749521	Hs.186545	ESTs	-	Sensitivity
127192	R08260	g760183	Hs.20131	ESTs	-	Sensitivity
128126	R09561	g761484	Hs.1369	decay accelerating factor for complement (CD55, Cromer blood group system)	DAF	Resistance
128126	R09672	g761595	Hs.1369	decay accelerating factor for complement (CD55, Cromer blood group system)	DAF	Resistance
153505	R48303	g810329	Hs.80552	dermatopontin	DPT	Resistance
153505	R48405	g810431	Hs.80552	dermatopontin	DPT	Resistance
154172	R52030	g813932	Hs.111732	IgG Fc binding protein	FC(GAMMA)BP	Sensitivity

Table 1

154654	R55185	g824480	Hs.3321	ESTs, Highly similar to IROQUOIS- CLASS HOMEODOMAIN PROTEIN IRX-3 [M.musculus]	-	Sensitivity
159455	H15746	g880566	Hs.74573	similar to vaccinia virus HindIII K4L ORF	HU-K4	Resistance
159455	H15747	g880567	Hs.74573	similar to vaccinia virus HindIII K4L ORF	HU-K4	Resistance
199243	R95869	g981529	Hs.35467	EST	-	Sensitivity
203348	H54285	g994432	Hs.9829	ESTs	-	Sensitivity
234907	H73080	g1046466	Hs.82007	KIAA0094 protein	KIAA0094	Resistance
234907	H73079	g1046465	Hs.82007	KIAA0094 protein	KIAA0094	Resistance
242642	H94977	g1102610	Hs.42041	EST	-	Sensitivity
243741	N49629	g1190795	Hs.44532	diubiquitin	UBD	Sensitivity
243741	N33920	g1154320	Hs.44532	diubiquitin	UBD	Sensitivity
245330	N76677	g1239255	Hs.251664	insulin-like growth factor 2 (somatomedin A)	IGF2	Resistance
245330	N54596	g1195916	Hs.251664	insulin-like growth factor 2 (somatomedin A)	IGF2	Resistance
261828	H99196	g1123864	Hs.226216	ESTs	-	Sensitivity
261828	N24479	g1138629	Hs.226216	ESTs	-	Sensitivity
274677	R84629	g943035	Hs.169338	ESTs	-	Resistance
274677	R85394	g943800	Hs.169338	ESTs	-	Resistance
277173	N44209	g1182737	Hs.181357	laminin receptor 1 (67kD, ribosomal protein SA)	LAMR1	Sensitivity
277173	N34316	g1155458	Hs.181357	laminin receptor 1 (67kD, ribosomal protein SA)	LAMR1	Sensitivity
284220	N53534	g1194700	Hs.171763	CD22 antigen	CD22	Resistance
291880	N67487	g1219612	Hs.83551	microfibrillar-associated protein 2	MFAP2	Resistance
291880	W03413	g1275326	Hs.83551	microfibrillar-associated protein 2	MFAP2	Resistance
295723	N66925	g1219050	Hs.49275	ESTs	-	Sensitivity
298417	N74131	g1231416	Hs.82961	trefoil factor 3 (intestinal)	TFF3	Sensitivity
322443	W39215	g1320924	Hs.238927	Homo sapiens mRNA; cDNA DKFZp434H1235 (from clone DKFZp434H1235); partial cds	-	Sensitivity

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322443	W16424	g1289598	Hs.238927	Homo sapiens mRNA; cDNA DKFZp434H1235 (from clone DKFZp434H1235); partial cds	-	Sensitivity
322723	W15465	g1289894	Hs.93231	ESTs	-	Sensitivity
322723	W39618	g1321460	Hs.93231	ESTs	-	Sensitivity
324699	W47134	g1331784	Hs.184019	Homo sapiens clone 23551 mRNA sequence	-	Sensitivity
324699	W47096	g1331890	Hs.184019	Homo sapiens clone 23551 mRNA sequence	-	Sensitivity
344942	W72861	g1383016	Hs.109299	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 3	PPFIA3	Sensitivity
344942	W75957	g1386331	Hs.109299	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 3	PPFIA3	Sensitivity
344958	W72892	g1383027	Hs.214507	ESTs	-	Sensitivity
344958	W76097	g1386341	Hs.214507	ESTs	-	Sensitivity
377799	AA777001	g2836332	Hs.79378	cyclin A1	CCNA1	Resistance
430077	AA010003	g1471050	Hs.79103	aminolevulinase, delta-, synthase 2 (sideroblastic/hypochromic anemia)	ALAS2	Sensitivity
430077	AA010004	g1471051	Hs.79103	aminolevulinase, delta-, synthase 2 (sideroblastic/hypochromic anemia)	ALAS2	Sensitivity
431944	AA678160	g2658682	Hs.117106	ESTs	-	Resistance
436348	AA776448	g2835782	Hs.122614	ESTs, Weakly similar to apoptotic protease activating factor 1 [M.musculus]	-	Sensitivity
436782	AA702821	g2705934	Hs.124778	ESTs	-	Sensitivity
451706	AA707650	g2717568	Hs.267289	polymerase (DNA directed), alpha	POLA	Sensitivity
460487	AA677706	g2658228	Hs.347	lactotransferrin	LTF	Sensitivity
461468	AA705029	g2714947	Hs.163036	ESTs	-	Resistance
488945	AA047077	g1524975	Hs.75733	amylase, alpha 2B; pancreatic	AMY2B	Sensitivity

Table 1

488945	AA047078	g1524976	Hs.180149	ESTs, Highly similar to ALPHA-AMYLASE 2B PRECURSOR [H. sapiens]	-	Sensitivity
489637	AA099445	g1645393	Hs.181060	apelin; peptide ligand for APJ receptor	APELIN	Sensitivity
489637	AA101878	g1645281	Hs.181060	apelin; peptide ligand for APJ receptor	APELIN	Sensitivity
490600	AA101616	g1648684	Hs.155210	FOS-like antigen 2	FOSL2	Resistance
490600	AA101617	g1648685	Hs.155210	FOS-like antigen 2	FOSL2	Resistance
506516	AA708619	g2718537	Hs.128856	CSR1 protein	CSR1	Resistance
506583	AA708512	g2718430	Hs.25537	cardiotrophin 1	CTF1	Resistance
510576	AA055768	g1548168	Hs.25615	YDD19 protein	YDD19	Sensitivity
510576	AA055880	g1548218	Hs.91011	anterior gradient 2 (Xenopus laevis) homolog	AGR2	Sensitivity
592540	AA160507	g1735874	Hs.195850	keratin 5 (epidermolysis bullosa simplex, Dowling-Meara/Kobner/Weber-Cockayne types)	KRT5	Sensitivity
592540	AA160595	g1735963	Hs.195850	keratin 5 (epidermolysis bullosa simplex, Dowling-Meara/Kobner/Weber-Cockayne types)	KRT5	Sensitivity
595037	AA173872	g1754021	Hs.194691	retinoic acid induced 3	RAI3	Sensitivity
595037	AA172400	g1751448	Hs.194691	retinoic acid induced 3	RAI3	Sensitivity
713263	AA283106	g1926031	Hs.89040	prepronociceptin	PNOC	Resistance
713263	AA283020	g1925944	Hs.89040	prepronociceptin	PNOC	Resistance
730871	AA417025	g2077124	Hs.98186	chromosome 21 open reading frame 22	C21ORF22	Sensitivity
731311	AA416767	g2077721	Hs.270266	ESTs, Weakly similar to ORF YKL201c [S. cerevisiae]	-	Resistance
739193	AA421218	g2100043	Hs.7678	cellular retinoic acid-binding protein 1	CRABP1	Sensitivity
739193	AA421217	g2100042	Hs.7678	cellular retinoic acid-binding protein 1	CRABP1	Sensitivity
741067	AA478436	g2207070	Hs.250581	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2	SMARCD2	Sensitivity
741067	AA402352	g2056264	Hs.250581	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2	SMARCD2	Sensitivity

Table 1

741891	AA402117	g2056108	Hs.170160	RAB2, member RAS oncogene family-like	RAB2L	Sensitivity
741891	AA401972	g2056031	Hs.170160	RAB2, member RAS oncogene family-like	RAB2L	Sensitivity
743465	AA609385	g2457813	Hs.112703	ESTs	-	Sensitivity
753587	AA478585	g2207219	Hs.167741	butyrophilin, subfamily 3, member A3	BTN3A3	Sensitivity
753587	AA479322	g2207878	Hs.167741	butyrophilin, subfamily 3, member A3	BTN3A3	Sensitivity
755881	AA496539	g2229860	Hs.179902	putative human HLA class II associated protein 1	PHAP1	Sensitivity
756463	AA481344	g2210896	Hs.8022	downregulated in renal cell carcinoma	TU3A	Sensitivity
756463	AA436401	g2141315	Hs.8022	downregulated in renal cell carcinoma	TU3A	Sensitivity
759163	AA442695	g2154573	Hs.118223	microfibrillar-associated protein 4	MFAP4	Resistance
759163	AA496022	g2229343	Hs.118223	microfibrillar-associated protein 4	MFAP4	Resistance
767993	AA418945	g2080755	Hs.29759	RNA POLYMERASE I AND TRANSCRIPT RELEASE FACTOR	PTRF	Resistance
767993	AA418829	g2080630	Hs.29759	RNA POLYMERASE I AND TRANSCRIPT RELEASE FACTOR	PTRF	Resistance
768602	AA425126	g2107197	Hs.98402	ESTs	-	Sensitivity
772446	AA405640	g2063132	Hs.105915	ESTs	-	Resistance
772446	AA405488	g2063071	Hs.105915	ESTs	-	Resistance
783729	AA446928	g2159593	Hs.173664	v-erb-b2 avian erythroblastic leukemia viral oncogene homolog 2 (neuro/glioblastoma derived oncogene homolog)	ERBB2	Sensitivity
783729	AA443351	g2156026	Hs.173664	v-erb-b2 avian erythroblastic leukemia viral oncogene homolog 2 (neuro/glioblastoma derived oncogene homolog)	ERBB2	Sensitivity
785847	AA449119	g2163139	Hs.200478	ubiquitin-conjugating enzyme E2M (homologous to yeast UBC12)	UBE2M	Resistance
788234	AA454080	g2167749	Hs.34853	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	ID4	Sensitivity

Table 1

788234	AA452493	g2166162	Hs. 34853	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	ID4	Sensitivity
788524	AA452937	g2166606	Hs. 99291	ESTs, Weakly similar to KIAA1006 protein [H.sapiens]	-	Resistance
788524	AA452801	g2166470	Hs. 99291	ESTs, Weakly similar to KIAA1006 protein [H.sapiens]	-	Resistance
788609	AA452899	g2166568	Hs. 213586	ESTs, Weakly similar to similar to KIAA0766 [H.sapiens]	-	Resistance
789369	AA464856	g2189740	Hs. 34853	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	ID4	Sensitivity
789369	AA453341	g2167010	Hs. 34853	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	ID4	Sensitivity
795378	AA453495	g2167164	Hs. 236463	Homo sapiens mRNA; cDNA DKFZp586I0521 (from clone DKFZp586I0521)	-	Sensitivity
809694	AA454702	g2177478	Hs. 7678	cellular retinoic acid-binding protein 1	CRABP1	Sensitivity
809694	AA456351	g2178927	Hs. 7678	cellular retinoic acid-binding protein 1	CRABP1	Sensitivity
809998	AA455195	g2177971	Hs. 274376	amylase, alpha 1A; salivary	AMY1A	Sensitivity
809998	AA454854	g2177630	Hs. 75733	amylase, alpha 2B; pancreatic	AMY2B	Sensitivity
810871	AA458981	g2183888	Hs. 171814	parathymosin	PTMS	Resistance
810871	AA459196	g2184103	Hs. 171814	parathymosin	PTMS	Resistance
814297	AA459105	g2184012	Hs. 73947	peptidase D	PEPD	Resistance
814297	AA459325	g2184232	Hs. 73947	peptidase D	PEPD	Resistance
815284	AA481608	g2211160	Hs. 73947	peptidase D	PEPD	Resistance
815284	AA481543	g2211095	Hs. 73947	peptidase D	PEPD	Resistance
837891	AA434092	g2139006	Hs. 271869	ESTs	-	Sensitivity
837891	AA434363	g2139277	Hs. 243010	ESTs, Moderately similar to GTP-BINDING PROTEIN TC10 [H.sapiens]	-	Sensitivity
841679	AA488699	g2218301	Hs. 10803	calcium and integrin binding protein (DNA-dependent protein kinase interacting protein)	SIP2-28	Sensitivity

Table 1 *continued*

841679	AA487575	g2217739	Hs. 10803	calcium and integrin binding protein (DNA-dependent protein kinase interacting protein)	SIP2-28	Sensitivity
842863	AA486403	g2216567	Hs. 75789	N-myc downstream regulated	NDRG1	Sensitivity
842863	AA489261	g2218863	Hs. 75789	N-myc downstream regulated	NDRG1	Sensitivity
845658	AA670144	g2631643	Hs. 61762	ESTs	-	Sensitivity
859586	AA668681	g2630180	Hs. 278736	cell division cycle 42 (GTP-binding protein, 25kD)	CDC42	Resistance
859761	AA668508	g2630007	Hs. 183986	poliovirus receptor-related 2 (herpesvirus entry mediator B)	PVRL2	Resistance
859858	AA679454	g2659976	Hs. 31132	steroidogenic acute regulatory protein	STAR	Resistance
897597	AA496846	g2230167	Hs. 278518	DKFZP434D174 protein	DKFZP434D174	Sensitivity
897597	AA496888	g2230209	Hs. 19614	gemin4	GEMIN4	Sensitivity
897641	AA496741	g2230062	Hs. 103804	heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	HNRPU	Sensitivity
897641	AA496792	g2230113	Hs. 139572	EST	-	Sensitivity
969769	AA772904	g2825746	Hs. 132884	heparan sulfate 6-O-sulfotransferase	HS6ST	Resistance
970649	AA774724	g2834058	Hs. 25615	YDD19 protein	YDD19	Resistance
1055543	AA620821	g2524760	Hs. 112911	EST	-	Sensitivity
1412238	AA844818	g2931269	Hs. 278399	amylase, alpha 2A; pancreatic	AMY2A	Sensitivity
1412245	AA844831	g2931282	Hs. 89717	carboxypeptidase A2 (pancreatic)	CPA2	Sensitivity
1456937	AA863449	g2955928	Hs. 1154	oviductal glycoprotein 1, 120kD	OVGP1	Resistance
1470220	AA865960	g2958236	Hs. 127286	ESTs	-	Resistance
1473682	AA916726	g3056118	Hs. 125262	DKFZP586G1624 protein	DKFZP586G1624	Sensitivity
1475842	AA872143	g2968321	Hs. 17820	Rho-associated, coiled-coil containing protein kinase 1	ROCK1	Resistance
1486028	AA912032	g3051424	Hs. 181059	ESTs	-	Resistance
1493205	AA878923	g2987888	Hs. 90680	ESTs, Weakly similar to WD40 protein C10orf1 [H.sapiens]	-	Resistance
1550776	AA912448	g3051840	Hs. 121529	ELK3, ETS-domain protein (SRF accessory protein 2)	ELK3	Resistance
1550909	AA913206	g3052598	Hs. 278606	G antigen 7	GAGE7	Resistance
1573946	AA938494	g3096622	Hs. 28555	programmed cell death 9	PDCD9	Resistance

Table 1

1574438	AA954935	g3118630	Hs.155324	matrix metalloproteinase 11 (stromelysin 3)	MMP11	Resistance
1588935	AA975612	g3151404	Hs.268557	pleckstrin homology-like domain, family A, member 3	PHLDA3	Resistance
1603560	AA996122	g3182611	Hs.73947	peptidase D	PEPD	Resistance
1605142	AA987928	g3173292	Hs.27457	ESTs	-	Resistance
1609538	AI000966	g3191520	Hs.104696	Homo sapiens mRNA for KIAA1324 protein, partial cds	-	Sensitivity
1623016	AI014781	g3229117	Hs.234903	EST	-	Sensitivity
1635203	AI003775	g3213285	Hs.127824	ESTs, Weakly similar to weak similarity to collagens [C.elegans]	-	Resistance
1635978	AI017801	g3232137	Hs.131201	ESTs	-	Resistance
1646649	AI025974	g3241587	Hs.131678	EST	-	Resistance
1916700	AI347629	g4084835	Hs.123107	kalikrein 1, renal/pancreas/salivary	KLK1	Resistance
1946534	AI351740	g4088946	Hs.890	lymphotoxin beta (TNF superfamily, member 3)	LTB	Sensitivity
1968246	AI285751	g3923984	Hs.118722	fucosyltransferase 8 (alpha (1,6) fucosyltransferase)	FUT8	Resistance

Table 2A

<u>EST</u>	<u>Sensitivity/Resistance</u>	<u>SEQ ID NO</u>
cohXres112c05	Sensitivity	1
jlhbab397f01	Sensitivity	18
jlhbab412e01	Sensitivity	16
jlhbab443e06	Resistance	2
jlhbab453e07	Sensitivity	17
jlhbac238e10	Resistance	3
jlhbad283g07	Resistance	19
jlhbae334b03	Sensitivity	4
jMhXp229h07	Resistance	15
jMhXp252a05	Resistance	5
johOf009h09	Resistance	6
johOf017b09	Resistance	7
johOf021e06	Resistance	8
johOf079g12	Resistance	9
johOf083h04	Resistance	10
johOf092b09	Resistance	11
johOf094e10	Resistance	12
jrhub001h03	Resistance	13
jrhoc127f11	Resistance	14

Table 2B

> cohXres112c05

AGGTACAAGCTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTCTACTGGAATCGTTNAAT
GNGTCTACTTNTCCACNCATAATTATAAAAGAATAAGAATCGACAAAAATATTTTNTTT
CCATAATATGTANAGGNGGTTGGTTTCTTTTTTTTTTTTTTCTTTCTTTAACTTTTT
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CG

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GACCATCTTTCTTTTGTGCTTTTTTTAAACATTTACTTTTCTTTAAGCCATAAGGATGC
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TANTAATAATGACTGGTCCGAATTTGGTTTTCGTTTTGTCTATTAAAGTCAATTTACTAA
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GGGCTCCACCTACAGTTGGTGGACATCCATGTGACAGAAATGGAGTCATCAGTTTATCA
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CTGGTGGAGCAAGTCTTTCTAGATAAGACCCCTGAATGCCTCATTCCATTGGCTGGGCTCC
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TCAAGAGTTATTTTTCTGACTGTCAAGTTTCAACATTCAAGTCTGTCCCCAACAGGCACC
ACACCGGGGTGGACTCCCTGTGTAACCTTCTCGCCACTGGCTCGGAGAGTAGACAGAGTTG
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CGGCTTTAGAACTAGTN

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AGAAAAATAACTCTTGATGCTGCTGTTTCGGAAGAGTTGGTTGAGCGCATCCTCAATATT
CCTTTTGTTCCTCTGGTAATTGGTGGTGCCCTGGCTGGGCTTTGTCTGGGAATATGGTAG
GTTGGTGATGGTGAAATTCAGGTAGAAGTGCTGGGTGCTGGAGCTGCTTGTGGTTGATA
AACTGATGACTCCATTTCTGTACATGGATGTCCACCAACTGGTAGGTGGAGCCCAGCCA
ATGGGAATGAGGCATTCAAGGTCTTATCTAGAAAGACTTGCTCCACCAGGCTGGGGTCCA
AATTGGAG

> johOf083h04

CCGCGGTGGCGGCCGCCCGGGCAGGTACATCACCTGCTGAGGGACATCCAGGACAAGGT
CACCACACTCTACAAAGGCAGTCAACTACATGACACATTCCGCTTCTGCCTGGTCACCAA
CTTGACGATGGACTCCGTGTGGTCACTGTCAAGGCATTGTTCTCCTCCAATTTGGACCC
CAGCCTGGTGGAGCAAGTCTTTCTAGATAAGACCCCTGAATGCCTCATTCCATTGGCTGGG
CTCCACCTACCAGTTGGTGGACATCCATGTGGCAGAAATGGAGTCATCAGTTTATCAACC
AACAAGCAGCTCCAGCACCCAGCACTTCTACCTGAATTTACCATCACCAACCTACCATA
TTCCCAGGACAAAGCCCAGCCAGGCACCACCAATTACCAGAGGAACAAAAGGAATATTGA
GGATGCGCTCAACCAACTCTTCCGAAACAGCAGCATCAAGAGT

> johOf092b09

ACCGCNGTGGCGGCCGCCCGGGCAGGTACATCACCTGCTGAGGGACTTTTNNGGACAAG
GTCACCACACTCTACAAAGGCAGTCAACTACATGACACATTCCGCTTCTGCCTGGTCACC
AATTGACGATGGACTCCGTGTTGGTCACTGTCAAGGCATTGTTCTCCTCCAATTTGGAC
CCCAGCCTGGTGGAGCAAGTCTTTCTAGATAAGACCCCTGAATGCCTCATTCCATTGGCTG
GGCTCCACCTACCAGTTGGTGGACATCCATGTGACAGAAATGGAGTCATCAGTTTATCA
AC

> johOf094e10

AGGGCGAATTGGAGCTCNCCGCGGTGGCGGCCGAGGTACCACCTGAAGGCCCTCACACTC
AACTTCACCATCTCCAATCTCCAGTATTCACCAGATATGGGCAAGGGCTCAGCTACATTC
AACTCCACCGAGGGGTCCTTCAGCACCTGCTCAGACCCTTGTTCAGAAGAGCAGCATG
GGCCCCCTTCTACTTGGGTTGCCAACTGATCTCCCTCAGGCCTGAGAAGGATGGGGCAGCC
ACTGGTGTGGACACCACCTGCACCTACCACCTGACCCTGTGGGCCCCGGGCTGGACATA
CAGCAGCTTTACTGGGAGCTGAGTCAGCTGACCCATGGGTGTACCCAACTGGGCTTCTA
TTGTCTGGACAGGATAGCCTCTTCATCAATGGCTATGCACCCCAAAATTTATCAATCC
GGGGCGAGGTACCTGCCCCGGGCGGGCCGCTTAAACTAGNGGGATCCCCNNGGCTTG

Table 2B

CAGGAATTTTCGATATTCAAGCTTATCGATACCCGTCCNACCTTCGAGGGGGGGG

> jrhub001h03

TGGGGGAAAGGGAGNNCCCAACGATCCTGGAACCTTTAANTNTGGAAAGAGTGAGATTTCAG
AAATCGCCACNACTGGACTTTAAGGGACGTCTGTGTGTCAGCACAANGGACTGGCACACAC
AGACACACNAGACCGANGANAACTGCANACAAATGGAGATACNAANACTTAGAAGGACA
GCTCCTTTACCTCATCTACTTGTCCAGAAGGTAAAAAGACACANCCAGAAAGAAAAGG
CATCNGCTCANCTCTCAGATCANGACANGCTGTGGATCTGTGGCGGTACT

> jrhub127f11

GCGTCCGAATTTCTGGGTACCCCGTATATAAGAAAATGTTAAAGTCAGGCAGGAAAAC
ATAGAATTTAAAGCCTTATAGTATATTATATAGNAAAGCCCTATATAGTATAGACAGAAAA
GTTTAGGGAAGGCCACAATTGCAAAGAAAAGTGGTGGTCACGGAACAAGGGAATGTCAT
ACAAATGTGGACACACACTGCGTTACTGAGCGCCACGTCTCATAGGTGAGAAGCATAACT
CTAGAAGGTGAGAAATGAGAATTTTCACTTCCATCCTTCCATTTGTTGTGTGACTCTGCC
ATTTACTTTCTTTNTTTTGTATTTTTCATTTTCTTTTAAAAATGGAAATATGAATTTTG
AATTTCTGCTCTATCTCACAGGTTTTTGTGGGGATGCATTTAAANGTTTAATTAGTAA
ATAATGGTAT

> jMhXp229h07

CCAAACTATTTGGACAGAATGGCTTCAAAGCTAGGCGNAAATGTTACATTATAAAAAAG
TTAAATATTACCTTCAATACCTGTCACTAGCCTACTGACAAATTATGACTAAACAAAGGT
ATTTGTATGACTATGTAATAGATCATCCGCTGAAAAGTAAAAACAAATAACAAAAAACT
TGTCCTAATGGGAAAGCATGCTTAATAAAAGGAAATGCACGAAGTTATAAACATGTTTTG
TNAGTAAGTATTCAGAATTTAAATATGTGATACATTTTATGATTGCTTAATGATCCTT
GGATGTCAGATTCCTTGGGTCTATTTATAGCTAAATTATAATGAAAAATTCAAGGCTTGC
TGNAGCAACTCTGTCAACAAATATATTAGTTTNGCTTATATATNTNGATTCTTTATGTGG
GAAAAATTACTACCC

> jlhab412e01

CGGCCGCCCCGGGGGATGCCGAGTCCCAAGAGGCCGAGTTTGAGAGGCTGGTGGCAGAAT
TCCCGGAGAAGGAGGCCAGCTGTCCCTGGTGGAAAGCGCAGGGCTGGCTGGTGATGGAGA
AGTCTTCTCCGGAGGGTGCTGCCGTGGTGCAGGAGGAGCTCAGGGAGCTGGCAGAGTCGT
GGCGGGCCTTGAGGCTGCTGGAAGAAAGTCTGCTGAGCCTCATCAGAACTGGCATCTGC
AGAGGATGGAAGTGGAATTCGGGGAAGAAAATGGTTTTACCAACAACATCCCAAAGTCAG
GATTTCTCATCAATCCCATGGATCCTATTTCCAGGCATCGTCGACGCGTGAGTCTGTCTA
GCAGGGCTGTGGGAGAAGGGGCCAGGCCCCAGGTCAAGAGGTGGGTAGGGGTCTCCAGCA
CAGGCCCTCCCTGTCTGGGGCAACATGCTCTGCTCTGAGGACTTGCCACGTCTCTGTCT
CATTTGAGCCTGC

> jlhab453e07

GCGTCCGGTTACAAAGTCAGGTTGTTATGGTTTGCATGACTTTGAGAAGCTAGTGGAATG
GAAATAAAGTTAGGAGCAGCAGGAGGAGGCTCTGTGTGGGCACATCTCCTTCAGGGGCAT
GGTACTGTTTCATGGACAGAGGAAGTCTATGGCATATGCTGGGACAGACAGTGAAGGGTA
GGTCTTACAAAGAGGCTTTACGTTAGAGTATAATAATCACTTATCTGTATGCATCTATGA
ATGATCTCACCGGATGTGAAGAATATGTATTTTAAAAACAGCATGAAACGGCCTGTAAT
CCCAGTACTTTTGGG

> jlhab397f01

ACTTATTGAATCATCGAATTCATTGAAGTTTGGCTCCAACCTATCATATCGCCGATGTTT
ACTTTTTCTTATTCTTATAAAGTTCTAAATTCAGAATGTGAGGTGGACAAATTCATTTT
AGTTCCACAAGTGGTAGCATTTAAATATCAGCAGCTTAAGTATTCAAATTAATAGATTG
CATTTTTTAAATGGTGAAATTCTGACAGTTTGCAGGGAAAAGGTGCTGAATATCTTGATA
TAATTTACATACTTCTATAAACAGGCATTTTTTATACCTTTGGAAAGATAAATGAGTAGAA
ACCAAGTATTTTACAATTTCTAATAGTTTATACTGACATGT

jlhab283g07

Table 2B

TCCGCTAAAAATTTGTTCTGGGCCTTTGGCTTAATTCAGAGATCTGCCCATGGGGTTCTATTACTTTGCTCTTTAACTTTG
TTCGATCCTTCTTGGATCAGTCTTGCAATTCATTCTTGTCTTTTCCTGAATAACATCTATGTTTTGCCCTCTTTTGAGTG
CTATCTTAATATGCCAGCCTATTTCTACCTTTCTTGTGCAGGGTAGCATAATTTTTACTTTCCATTATACCTCAGTCCCA
CACCTTGTGTCTGTTTATTTCAATACCTAAGATACTTATCCTCAGTTCCTAGCTTACTTTAGTTCTGAAAGTTGGATAT
CCATAATTGTAGTGGTCTTAAATCTGTAAAACACATATGGATGGGAAACCACTGAATAATGTAAATAAATATGAATAACG
ATGATAAAATAAAAATGATAAAAATAACTGAGTTCAATGATATTAAAAACATAAGTCA

Table 3A

Clone	Annotation
242642	EST//(Hs.42041;)
121661	ESTs//(Hs.226410;)
32567	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
788234	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
789369	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
27544	prominin (mouse)-like 1//(Hs.112360;NM_006017)
322723	ESTs//(Hs.93231;)
243741	diubiquitin//(Hs.44532;NM_006398)
1055543	EST//(Hs.112911;)
245330	insulin-like growth factor 2 (somatomedin A)//(Hs.251664;NM_000612)
1574438	matrix metalloproteinase 11 (stromelysin 3)//(Hs.155324;NM_005940)
234907	KIAA0094 protein//(Hs.82007;)
1456937	oviductal glycoprotein 1, 120kD//(Hs.1154;NM_002557)
1493205	ESTs, Weakly similar to WD40 protein Ciao 1 [H.sapiens]//(Hs.90680;)
506583	cardiotrophin 1//(Hs.25537;NM_001330)
jlhbac238e10	
970649	ESTs//(Hs.116561;)
291880	microfibrillar-associated protein 2//(Hs.83551;NM_002403)
731311	ESTs, Weakly similar to ORF YKL201c [S.cerevisiae]//(Hs.270266;)
1588935	ESTs, Highly similar to TDAG51/lpl homologue 1 [H.sapiens]//(Hs.110222;)
859586	cell division cycle 42 (GTP-binding protein, 25kD)//(Hs.146409;NM_001791)
1550909	G antigen 7//(Hs.184794;)
810871	parathymosin//(Hs.171814;NM_002824)
159455	similar to vaccinia virus HindIII K4L ORF//(Hs.74573;)
jlhbad283g07	
814297	peptidase D//(Hs.73947;NM_000285)
1635978	ESTs//(Hs.131201;)
785847	ubiquitin-conjugating enzyme E2M (homologous to yeast UBC12)//(Hs.200478;NM_003969)
274677	ESTs//(Hs.169338;)
859761	Untitled//(Hs.270880;)
377799	cyclin A1//(Hs.79378;NM_003914)
713263	prepronociceptin//(Hs.89040;)
1916700	kallikrein 1, renal/pancreas/salivary//(Hs.123107;NM_002257)
490600	FOS-like antigen 2//(Hs.155210;NM_005253)

Table 3B

<u>Clone</u>	<u>Annotation</u>
1916700	kallikrein 1, renal/pancreas/salivary/(Hs.123107;NM_002257)
490600	FOS-like antigen 2/(Hs.155210;NM_005253)
johOf021e06	
859761	Untitled/(Hs.270880;)
814297	peptidase D/(Hs.73947;NM_000285)
johOf009h09	
1605142	ESTs/(Hs.27457;)
johOf017b09	
johOf092b09	
johOf083h04	
451706	polymerase (DNA directed), alpha/(Hs.267289;)
506583	cardiotrophin 1/(Hs.25537;NM_001330)
753587	butyrophilin, subfamily 3, member A3/(Hs.167741;NM_006994)
789369	inhibitor of DNA binding 4, dominant negative helix-loop- helix protein/(Hs.34853;NM_001546)
261828	UNIGENE-ambiguity: Hs.226216::Hs.270258! ESTs/(Hs.226216;)
969769	heparan sulfate 6-O- sulfotransferase/(Hs.132884;NM_004807)
1603560	peptidase D/(Hs.73947;NM_000285)
32567	inhibitor of DNA binding 4, dominant negative helix-loop- helix protein/(Hs.34853;NM_001546)
772446	ESTs/(Hs.105915;)
159455	similar to vaccinia virus HindIII K4L ORF/(Hs.74573;)
788234	inhibitor of DNA binding 4, dominant negative helix-loop- helix protein/(Hs.34853;NM_001546)
johOf094e10	
897597	DKFZP434B131 protein/(Hs.19614;)
810871	parathymosin/(Hs.171814;NM_002824)
731311	ESTs, Weakly similar to ORF YKL201c [S.cerevisiae]/(Hs.270266;)
753587	butyrophilin, subfamily 3, member A3/(Hs.167741;NM_006994)
322443	ESTs/(Hs.238927;)
66498	ESTs/(Hs.21688;)
242642	EST/(Hs.42041;)
970649	ESTs/(Hs.116561;)
841679	calcium and integrin binding protein (DNA-dependent protein kinase interacting protein)/(Hs.10803;)
1588935	ESTs, Highly similar to TDAG51/lpl homologue 1 [H.sapiens]/(Hs.110222;)
785847	ubiquitin-conjugating enzyme E2M (homologous to yeast UBC12)/(Hs.200478;NM_003969)
johOf079g12	
121661	ESTs/(Hs.226410;)

Table 4

Clone	Annotation
66420	ESTs/(Hs.189713;)
460487	lactotransferrin/(Hs.347;NM_002343)
cohXres112c05	
242642	EST/(Hs.42041;)
430077	aminolevulinate, delta-, synthase 2 (sideroblastic/hypochromic anemia)/(Hs.79103;NM_000032)
1623016	EST/(Hs.234903;)
27544	prominin (mouse)-like 1/(Hs.112360;NM_006017)
121661	ESTs/(Hs.226410;)
324699	Homo sapiens clone 23551 mRNA sequence/(Hs.184019;)
199243	EST/(Hs.35467;)
741067	UNIGENE-ambiguity: Hs.250581::Hs.69469! SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2/(Hs.250581;NM_003077)
127192	ESTs/(Hs.20131;)
1055543	EST/(Hs.112911;)
344942	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 3/(Hs.109299;)
154172	IgG Fc binding protein/(Hs.111732;NM_003890)
756463	downregulated in renal cell carcinoma/(Hs.8022;NM_007177)
277173	laminin receptor 1 (67kD, ribosomal protein SA)/(Hs.181357;NM_002295)
489637	ESTs/(Hs.22793;)
743465	ESTs/(Hs.112703;)
755881	ESTs/(Hs.179902;)
595037	retinoic acid induced 3/(Hs.194691;NM_003979)
66317	H1 histone family, member 2/(Hs.7644;NM_005319)
795378	ESTs/(Hs.236463;)
1035784	
344958	ESTs/(Hs.214507;)
436782	ESTs/(Hs.124778;)
730871	ESTs/(Hs.98186;)
298417	trefoil factor 3 (intestinal)/(Hs.82961;NM_003226)
845658	ESTs/(Hs.61762;)
510576	UNIGENE-ambiguity: Hs.122576::Hs.91011! ESTs/(Hs.122576;)
203348	ESTs/(Hs.9829;)
783729	v-erb-b2 avian erythroblastic leukemia viral oncogene homolog 2 (neuro/glioblastoma derived oncogene homolog)/(Hs.173664;NM_004448)
241120	
768602	ESTs/(Hs.98402;)
51221	ortholog of rat pippin/(Hs.106635;)
897641	UNIGENE-ambiguity: Hs.103804::Hs.139572! heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)/(Hs.103804;NM_004501)
436348	ESTs, Weakly similar to apoptotic protease activating factor 1 [M.musculus]/(Hs.122614;)
295723	ESTs/(Hs.49275;)

Table 4

592540	keratin 5 (epidermolysis bullosa simplex, Dowling-Meara/Kobner/Weber-Cockayne types)/(Hs.195850;NM_000424)
970649	ESTs/(Hs.116561;)
1456937	oviductal glycoprotein 1, 120kD/(Hs.1154;NM_002557)
1493205	ESTs, Weakly similar to WD40 protein Ciao 1 [H.sapiens]/(Hs.90680;)
1573946	ESTs, Weakly similar to B0511.8 [C.elegans]/(Hs.28555;)
1470220	ESTs/(Hs.127286;)
785847	ubiquitin-conjugating enzyme E2M (homologous to yeast UBC12)/(Hs.200478;NM_003969)
461468	ESTs/(Hs.163036;)
33267	insulin-like growth factor binding protein 2 (36kD)/(Hs.162;)
506516	ESTs/(Hs.120152;)
1486028	ESTs/(Hs.181059;)
1550776	ELK3, ETS-domain protein (SRF accessory protein 2) NOTE: Symbol and name provisional./(Hs.121529;NM_005230)
154929	
1475842	Rho-associated, coiled-coil containing protein kinase 1/(Hs.17820;NM_005406)
26617	activated leucocyte cell adhesion molecule/(Hs.10247;NM_001627)
jMhXp252a05	
1646649	EST/(Hs.131678;)
1968246	fucosyltransferase 8 (alpha (1,6) fucosyltransferase)/(Hs.118722;NM_004480)
153505	dermatopontin/(Hs.80552;NM_001937)
jlhbab443e06	
969769	heparan sulfate 6-O-sulfotransferase/(Hs.132884;NM_004807)
1635203	ESTs, Weakly similar to weak similarity to collagens [C.elegans]/(Hs.127824;)
814297	peptidase D/(Hs.73947;NM_000285)
1916700	kallikrein 1, renal/pancreas/salivary/(Hs.123107;NM_002257)
859858	steroidogenic acute regulatory protein/(Hs.3132;NM_000349)
810871	parathymosin/(Hs.171814;NM_002824)

Table 5

Clone	Annotation
242642	EST//(Hs.42041;)
121661	ESTs//(Hs.226410;)
739193	cellular retinoic acid-binding protein 1//(Hs.7678;NM_004378)
32567	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
788234	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
809694	cellular retinoic acid-binding protein 1//(Hs.7678;NM_004378)
27544	prominin (mouse)-like 1//(Hs.112360;NM_006017)
809998	UNIGENE-ambiguity: Hs.252475::Hs.250817! amylase, alpha 2B; pancreatic//(Hs.252475;)
789369	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
1412238	amylase, alpha 2A; pancreatic//(Hs.75733;NM_000699)
755881	ESTs//(Hs.179902;)
743465	ESTs//(Hs.112703;)
460487	lactotransferrin//(Hs.347;NM_002343)
1609538	ESTs//(Hs.104696;)
298417	trefoil factor 3 (intestinal)//(Hs.82961;NM_003226)
1412245	carboxypeptidase A2 (pancreatic)//(Hs.89717;NM_001869)
322723	ESTs//(Hs.93231;)
jlhbab397f01	
jlhbae334b03	
154654	ESTs, Highly similar to IROQUOIS-CLASS HOMEODOMAIN PROTEIN IRX-3 [M.musculus]//(Hs.3321;)
842863	N-myc downstream regulated//((Hs.75789;NM_006096)
1473682	DKFZP586G1624 protein//((Hs.125262;)
285507	EST//(Hs.161495;)
277173	laminin receptor 1 (67kD, ribosomal protein SA)//(Hs.181357;NM_002295)
jlhbab453e07	
488945	UNIGENE-ambiguity: Hs.252475::Hs.180149! amylase, alpha 2B; pancreatic//(Hs.252475;)
jlhbab412e01	
741891	RAB2, member RAS oncogene family- like//((Hs.170160;NM_004761)
122906	ESTs//(Hs.186545;)
1946534	lymphotoxin beta (TNF superfamily, member 3)//(Hs.890;NM_002341)
837891	UNIGENE-ambiguity: Hs.271869::Hs.267654! ESTs//(Hs.271869;)
1493205	ESTs, Weakly similar to WD40 protein Ciao 1 [H.sapiens]//(Hs.90680;)
jrhoc127f11	
jlhbac238e10	
713263	prepronociceptin//((Hs.89040;)
970649	ESTs//(Hs.116561;)
jlhbad283g07	
767993	Homo sapiens mRNA; cDNA DKFZp586L2123 (from clone DKFZp586L2123)//(Hs.29759;)

Table 5

1456937	oviductal glycoprotein 1, 120kD//(Hs.1154;NM_002557)
284220	CD22 antigen//(Hs.171763;NM_001771)
1635978	ESTs//(Hs.131201;)
731311	ESTs, Weakly similar to ORF YKL201c [S.cerevisiae]//(Hs.270266;)
jMhXp229h07	
785847	ubiquitin-conjugating enzyme E2M (homologous to yeast UBC12) //(Hs.200478;NM_003969)
788524	ESTs, Weakly similar to KIAA1006 protein [H.sapiens] //(Hs.99291;)
759163	microfibrillar-associated protein 4 //(Hs.118223;)
788609	ESTs, Weakly similar to similar to KIAA0766 [H.sapiens] //(Hs.213586;)
128126	decay accelerating factor for complement (CD55, Cromer blood group system) //(Hs.1369;)
1635203	ESTs, Weakly similar to weak similarity to collagens [C.elegans] //(Hs.127824;)
859858	steroidogenic acute regulatory protein //(Hs.3132;NM_000349)
jrhob001h03	
431944	ESTs //(Hs.117106;)
815284	peptidase D //(Hs.73947;NM_000285)

Table 6

<u>Clone</u>	<u>Annotation</u>
242642	EST//(Hs.42041;)
32567	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
788234	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
789369	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein//(Hs.34853;NM_001546)
814297	peptidase D//(Hs.73947;NM_000285)
859761	Untitled//(Hs.270880;)
1916700	kallikrein 1, renal/pancreas/salivary//(Hs.123107;NM_002257)
490600	FOS-like antigen 2//(Hs.155210;NM_005253)